SEQUENCE LISTING

<110> ADViSYS, Inc.

<120> Insulin-Like Growth Factor ("IGF-I") Plasmid Mediated
Supplementation for Therapeutic Applications

<130> 108328.00172 - AVSI-0034

<160> 9

<170> PatentIn version 3.1

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<212> DNA
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<223> Nucleic acid sequence for the pAV2001 plasmid.

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<212>   DNA
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<220>
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<223>  This is a synthetic promoter SPc5-12

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<210>  4
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<211> 153
<212> PRT
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<220>
<223> This is an IGF-I amino acid sequence.

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Phe Tyr Leu Ala Leu Cys Leu Leu Thr Phe Thr Ser Ser Ala Thr Ala
            35   40   45

Gly Pro Glu Thr Leu Cys Gly Ala Glu Leu Val Asp Ala Leu Gln Phe
    50   55   60

Val Cys Gly Asp Arg Gly Phe Tyr Phe Asn Lys Pro Thr Gly Tyr Gly
65   70   75   80

Ser Ser Ser Arg Arg Ala Pro Gln Thr Gly Ile Val Asp Glu Cys Cys
                85   90   95

Phe Arg Ser Cys Asp Leu Arg Arg Leu Glu Met Tyr Cys Ala Pro Leu
            100   105   110

Lys Pro Ala Lys Ser Ala Arg Ser Val Arg Ala Gln Arg His Thr Asp
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Met Pro Lys Thr Gln Lys Glu Val His Leu Lys Asn Ala Ser Arg Gly
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Ser Ala Gly Asn Lys Asn Tyr Arg Met
145   150

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<211> 2237
<212> DNA
<213> artificial sequence

<220>
<223> Nucleotide sequence for the skeletal alpha actin 3' end.

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<210>  6
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<220>
<223>  Nucleic acid sequence of a human growth hormone 3' UTR.

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<210>  7
<211>  215
<212>  PRT
<213>  Artificial Sequence

<220>
<223>  Protein sequence of vascular endothelial growth factor

<400>  7

Met Asn Phe Leu Leu Ser Trp Val His Trp Ser Leu Ala Leu Leu Leu
1               5                   10                  15


Tyr Leu His His Ala Lys Trp Ser Gln Ala Ala Pro Met Ala Glu Gly
            20                  25                  30


Gly Gly Gln Asn His His Glu Val Val Lys Phe Met Asp Val Tyr Gln
        35                  40                  45


Arg Ser Tyr Cys His Pro Ile Glu Thr Leu Val Asp Ile Phe Gln Glu
    50                  55                  60


Tyr Pro Asp Glu Ile Glu Tyr Ile Phe Lys Pro Ser Cys Val Pro Leu
65                  70                  75                  80
```

```
Met Arg Cys Gly Gly Cys Cys Asn Asp Glu Gly Leu Glu Cys Val Pro
              85                    90                    95


Thr Glu Glu Ser Asn Ile Thr Met Gln Ile Met Arg Ile Lys Pro His
             100                 105                 110


Gln Gly Gln His Ile Gly Glu Met Ser Phe Leu Gln His Asn Lys Cys
             115                 120                 125


Glu Cys Arg Pro Lys Lys Asp Arg Ala Arg Gln Glu Lys Lys Ser Val
     130             135                 140


Arg Gly Lys Gly Lys Gly Gln Lys Arg Lys Arg Lys Lys Ser Arg Tyr
145             150                 155                     160


Lys Ser Trp Ser Val Pro Cys Gly Pro Cys Ser Glu Arg Arg Lys His
                 165             170                     175


Leu Phe Val Gln Asp Pro Gln Thr Cys Lys Cys Ser Cys Lys Asn Thr
                 180                 185                 190


Asp Ser Arg Cys Lys Ala Arg Gln Leu Glu Leu Asn Glu Arg Thr Cys
             195                 200                 205


Arg Cys Asp Lys Pro Arg Arg
     210                 215


<210>   8
<211>   782
<212>   DNA
<213>   artificial sequence

<220>
<223>   Nucleic acid sequence of a plasmid pUC-18 origin of replicaiton

<400>   8
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tcagctcact caaaggcggt aatacggtta tccacagaat caggggataa cgcaggaaag     120

aacatgtgag caaaaggcca gcaaaaggcc aggaaccgta aaaaggccgc gttgctggcg     180

ttttttccata ggctccgccc ccctgacgag catcacaaaa atcgacgctc aagtcagagg     240

tggcgaaacc cgacaggact ataaagatac caggcgtttc cccctggaag ctccctcgtg     300

cgctctcctg ttccgaccct gccgcttacc ggatacctgt ccgcctttct cccttcggga     360

agcgtggcgc tttctcatag ctcacgctgt aggtatctca gttcggtgta ggtcgttcgc     420

tccaagctgg gctgtgtgca cgaaccccccc gttcagcccg accgctgcgc cttatccggt     480
```

```
aactatcgtc ttgagtccaa cccggtaaga cacgacttat cgccactggc agcagccact      540

ggtaacagga ttagcagagc gaggtatgta ggcggtgcta cagagttctt gaagtggtgg      600

cctaactacg gctacactag aaggacagta tttggtatct gcgctctgct gaagccagtt      660

accttcggaa aaagagttgg tagctcttga tccggcaaac aaaccaccgc tggtagcggt      720

ggtttttttg tttgcaagca gcagattacg cgcagaaaaa aaggatctca agaagatcct      780

tt                                                                     782


<210>   9
<211>   5
<212>   DNA
<213>   artificial sequence

<220>
<223>   This is a NEO ribosomal binding site

<400>   9
tcctc                                                                    5
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